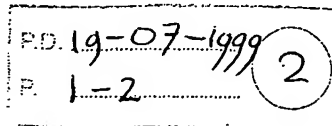


ID AQ743305 standard; DNA; GSS; 1141 BP.

XX
AC AQ743305;XX
SV AQ743305.1

XP-002214818

XX
DT 19-JUL-1999 (Rel. 60, Created)XX
DT 04-MAR-2000 (Rel. 63, Last updated, Version 2)XX
DE HS_5387_B2_B02_SP6 RPCI-11 Human Male BAC Library Homo sapiens genomic
DE clone Plate=963 Col=4 Row=D, genomic survey sequence.XX
KW GSS.

XX

OS Homo sapiens (human)

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;

OC Eutheria; Primates; Catarrhini; Hominidae; Homo.

XX

RN [1]

RP 1-1141

RX MEDLINE; 99380589.

RA Mahairas G.G., Wallace J.C., Smith K., Swartzell S., Holzman T., Keller A.,

RA Shaker R., Furlong J., Young J., Zhao S., Adams M.D., Hood L.;

RT "Sequence-tagged connectors: A sequence approach to mapping and scanning

RT the human genome";

RL Proc. Natl. Acad. Sci. U.S.A. 96(17):9739-9744(1999).

XX

CC Contact: Mahairas GG, Wallace JC, Hood L

CC High Throughput Sequencing Center

CC University of Washington

CC 401 Queen Anne Avenue North, Seattle, WA 98109, USA

CC Tel: (206) 616-3618

CC Fax: (206) 616-3887

CC Email: jwallace@u.washington.edu

CC Clones are derived from the human BAC library RPCI-11. For BAC

CC library availability, please contact Pieter de Jong

CC (pieter@dejong.med.buffalo.edu). Clones may be purchased from

CC BACPAC Resources (http://bacpac.med.buffalo.edu/ordering_bac.htm)

CC or from Resear h Genetics (info@resgen.com). BAC end Web Server:

CC <http://www.htsc.washington.edu>

CC Plate: 963 row: D column: 4

CC Seq primer: SP6

CC Class: BAC ends

CC High quality sequence stop: 1141.

XX

FH Key Location/Qualifiers

FH

FT source 1..1141

FT /db_xref="taxon:9606"

FT /note="Vector: pBACe3.6; Genomic sequence of BAC ends"

FT /sex="male"

FT /organism="Homo sapiens"

FT /clone="Plate=963 Col=4 Row=D"

FT /clone_lib="RPCI-11 Human Male BAC Library"

XX

SQ Sequence 1141 BP; 760 A; 301 C; 16 G; 15 T; 49 other;

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attnnaaggg cggccaanaa anccctttan aaggaaaaan tacanggacc ncnnaaaana	120
nannaaaaaa aaaaaacaaa caaaaaatta aaacaaaaaa acaaaaaaaa ccacaaaaaa	180
cacacacaa naaaacacaa aaaacaaaaa acaaaaccac aacaacacaa caaaaaacca	240
aaacaaacaa aaaaaacaaa aaaaaaaaac aaaaaaaaac aaaaaaaaac ccanacacac	300
aacaacacaa caacaacacc aaaacacaca cacaacacnc accaaaaaaa caacaacaca	360
aaacacaaca aanacaaac aaaccacgac accaaacaac aacacacaca aaacacaac	420
aaaaaacaaa aaaaaccaac aaacaaccaa aaaaaaaaac caaaaccaac aacacacaaa	480

caacaccaac	aacaacaaaa	caacaaaaca	aaaacaacaa	caaaccacaa	aaaaaacaaa	540
cacacaacaa	ncacaaaaaa	acaacaaaac	aaaacaacaa	aaaaaacaaa	acaaaaaaan	600
acaaaaacaa	caaacacac	aaccacacaa	aacaacacaa	acaacacaa	aacaacacca	660
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a						1141

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